

#2

OIPe

RAW SEQUENCE LISTING

DATE: 12/21/2001

PATENT APPLICATION: US/10/010,940

TIME: 08:56:33

Input Set : A:\427d3.app.txt

Output Set: N:\CRF3\12212001\J010940.raw

P.5

ENTERED

4 <110> APPLICANT: Xu, Jiangchun
 5 Dillon, Davin C.
 6 Mitcham, Jennifer L.
 7 Harlocker, Susan Louise
 8 Jiang Yuqui
 9 Reed, Steven G.
 10 Kalos, Michael
 11 Fanger, Gary
 12 Retter, Mark
 13 Solk, John
 14 Day, Craig
 16 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
 17 DIAGNOSIS OF PROSTATE CANCER
 19 <130> FILE REFERENCE: 210121.427D3
 C--> 21 <140> CURRENT APPLICATION NUMBER: US/10/010,940
 22 <141> CURRENT FILING DATE: 2001-12-05
 24 <160> NUMBER OF SEQ ID NOS: 575
 26 <170> SOFTWARE: FastSEQ for Windows Version 3.0
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 29 <211> LENGTH: 814
 30 <212> TYPE: DNA
 31 <213> ORGANISM: Homo sapien
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 34 <221> NAME/KEY: misc_feature
 35 <222> LOCATION: (1)...(814)
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 40 atcaaactcg aggggtgtct ggaggacttc aatacacctc ccccatagtg gaatcagctt 120
 41 ccagggggtc cagtcctctt cttacttcca tccccatccc atgccaaagg aagaccctcc 180
 42 ctccttggtc cacagccttc tctaggettcc ctagtgcttc caggacagag tgggttatgt 240
 43 tttcagctcc atccttgcctg tgagtgtctg gtgcgttggt cctccagctt ctgctcagtg 300
 44 cttcatggac agtgtccagc acatgtcaact ctccactctc tcagtgtgga tccactagtt 360
 45 ctagagcggc cgccaccgcg gtggagctcc agcttttggt cccttttagtg aggggttaatt 420
 46 gcgcgcttgg cgtaaatcatg gtcataactg tttcctgtgt gaaattgtta tccgctcaca 480
 47 attccacaca acatacgagc cgggaagcata aagtgtaaaag cctgggggtgc ctaatgagtg 540
 W--> 48 anctaactca cattaattgc gttgcgctca ctgncgctt tccagtcngg aaaactgtcg 600
 W--> 49 tgccagctgc attaatgaat cggccaaacgc ncgggggaaaa gcgggtttgcg ttttgggggc 660
 W--> 50 tcttccgctt ctcgctcaact nantcctgcg ctcggtcntt cggtctcggg gaacgggtatc 720
 W--> 51 actcctcaaa ggnggtatta cggttatccn naaatcnggg gataccngg aaaaaanttt 780
 W--> 52 aacaaaaggg cancaaaggg cngaaacgta aaaa 814
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 55 <211> LENGTH: 816
 56 <212> TYPE: DNA
 57 <213> ORGANISM: Homo sapien
 59 <220> FEATURE:
 60 <221> NAME/KEY: misc_feature

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66 ttcattggtg ttggagcaat agaaccacag ttctacgagc tgctgatcaa aggacttgga      120
67 cttaaagtctg atgaacttcc caatcagatg agcatggatg attggccaga aatgaagaag      180
68 aagttttgcag atgtatttgc aaagaagacg aaggcagagt ggtgtcaa atttgacggc      240
69 acagatgcct gtgtgactcc ggttctgact tttagaggag ttgttcatca tgatcacaac      300
70 aaggaacggg gctcgtttat caccagttag gaggcaggag tgagcccccg ccctgcacct      360
71 ctgctgttaa acacccacg catcccttct ttcaaaaggg atccactagt tctagaagcg      420
72 gccgccaccg cgggtggagct ccagcttttg ttccctttag tgagggttaa ttgcgcgctt      480
73 ggcgtaatac tggatcatag tgtttcctgt gtgaaattgt tatccgctca caattcccc      540
W--> 74 aacatacgag ccggaacata aagtgttaag cctgggggtgc ctaatgantg agctaactcn      600
W--> 75 cattaattgc gttgcgctca ctgcccgctt tccagtcggg aaaactgtcg tgccactgcn      660
W--> 76 ttantgaatc ngccaccccc cgggaaaagg cggttgcntt ttgggcctct tccgctttcc      720
W--> 77 tcgctcattg atcctngenc ccggtcttcg gctgcggnga acggttcaact cctcaaaggc      780
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81 <211> LENGTH: 773
82 <212> TYPE: DNA
83 <213> ORGANISM: Homo sapien
85 <220> FEATURE:
86 <221> NAME/KEY: misc_feature
87 <222> LOCATION: (1)...(773)
88 <223> OTHER INFORMATION: n = A,T,C or G
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92 tcctgctcct cactggtgat aaacgagccc cgttccttgt tgtgatcatg atgaacaacc      120
93 tcctcaaaaag tcagaaccgg agtcacacag gcatctgtgc cgtcaaagat ttgacaccac      180
94 totgccttcg tcttctttgc aaatacatct gcaaaacttct tcttcatttc tggccaatca      240
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96 tcgtagaact ggggttctat tgctccaaca gccatgaatt ccccatctgc tgtcctgtaa      360
97 gtcgtataga aagtgctcc accatccaac atgttctgtc ctcgaggggg ggcccgttac      420
W--> 98 ccaattcgcc ctatantgag tcgtattacg cgcgctcact ggccgtcgtt ttacaacgtc      480
99 gtgactggga aaaccctggg cgttaccac ttaatgcct tgcagcacat cccctttc      540
W--> 100 ccagctgggc gtaatanca aaaggccgc accgatcgcc cttccaacag ttgcgcacct      600
W--> 101 gaatgggnaa atgggacccc cctgttaccg cgcattnaac ccccgnggg tttngttgtt      660
W--> 102 acccccacnt nnaccgctta cactttgcca gcgccttanc gcccgctccc tttnccttt      720
W--> 103 cttecccttc tttcncccn ctttcccccg gggtttcccc cntcaaacc cna      773
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106 <211> LENGTH: 828
107 <212> TYPE: DNA
108 <213> ORGANISM: Homo sapien
110 <220> FEATURE:
111 <221> NAME/KEY: misc_feature
112 <222> LOCATION: (1)...(828)
113 <223> OTHER INFORMATION: n = A,T,C or G
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117 aatgggcaga cacaggtgta tgccaatggt tctgaaatgg gtataatttc gtcctctcct 120
118 tcggaacact ggctgtctct gaagacttct cgctcagttt cagtgaggac acacacaaag 180
119 acgtgggtga ccatgttggt tgtgggggtgc agagatggga ggggtggggc ccaccctgga 240
120 agagtggaca gtgacacaag gtggacactc tctacagatc actgaggata agctggagcc 300
W--> 121 acaatgcatg aggcacacac acagcaagga tgacnctgta aacatagccc acgctgtcct 360
W--> 122 gngggcactg ggaagcctan atnaggccgt gagcanaaag aaggggagga tccactagtt 420
W--> 123 ctanagcggc cgccaccgcg gtgganctcc ancttttggt cccttttagtg agggttaatt 480
W--> 124 gcgcgcttgg cntaatcatg gtcatanctn tttcctgtgt gaaattgtta tccgctcaca 540
W--> 125 attccacaca acatacganc cggaacata aantgtaaac ctgggggtgcc taatgantga 600
W--> 126 ctaactcaca ttaattgcgt tgcgctcact gcccgctttc caatcnggaa acctgtcttg 660
W--> 127 ccncttgcat tnatgaatcn gccaaccccc ggggaaaagc gtttgcggtt tgggcgctct 720
W--> 128 tccgcttctc cnetcantta ntccctncnc tccgctcatt cggtgcngc aaaccgggtc 780
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133 <212> TYPE: DNA
134 <213> ORGANISM: Homo sapien
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137 <221> NAME/KEY: misc_feature
138 <222> LOCATION: (1)...(834)
139 <223> OTHER INFORMATION: n = A,T,C or G
141 <400> SEQUENCE: 5
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143 agtttttaatt gcatccaaag tactaacaaa aactctagca atcaagaatg gcagcatggt 120
144 atttttataac aatcaacacc tgtggctttt aaaatttggt tttcataaga taattttatac 180
145 tgaagtaaat ctagccatgc ttttaaaaaa tgctttaggt cactccaagc ttggcagtta 240
146 acatttgcca taaacaataa taaaacaatc acaatttaat aaataacaaa tacaacattg 300
147 taggccataa tcatatacag tataaggaaa aggtggtagt gttgagtaag cagttattag 360
148 aatagaatac cttggcctct atgcaaatat gtctagacac tttgattcac tcagccctga 420
149 cattcagttt tcaaagtagg agacaggttc tacagtatca ttttacagtt tccaacacat 480
150 tgaaaacaag tagaaaatga tgagttgatt tttattaatg cattacatcc tcaagagtta 540
151 tcaccaaccc ctacagttata aaaaattttc aagttatatt agtcatataa cttggtgtgc 600
152 ttatttttaa ttagtgctaa atggattaag tgaagacaac aatgggtccc taatgtgatt 660
W--> 153 gatattggtc atttttacca gcttctaaat ctnaactttc aggcctttga actggaacat 720
W--> 154 tgnatnacag tgttccanag ttncaaccta ctggaacatt acagtgtgct tgattcaaaa 780
W--> 155 tgttattttg ttaaaaatta aattttaacc tgggtggaaa ataatttgaa atna 834
157 <210> SEQ ID NO: 6
158 <211> LENGTH: 818
159 <212> TYPE: DNA
160 <213> ORGANISM: Homo sapien
162 <220> FEATURE:
163 <221> NAME/KEY: misc_feature
164 <222> LOCATION: (1)...(818)
165 <223> OTHER INFORMATION: n = A,T,C or G
167 <400> SEQUENCE: 6
168 tttttttttt tttttttttt aagaccctca tcaatagatg gagacataca gaaatagtca 60
169 aaccacatct acaaaatgcc agtatcaggc ggcggcttcg aagccaaagt gatgtttgga 120
170 tgtaaagtga aatattagtt ggcggatgaa gcagatagtg aggaaagtgt agccaataat 180
171 gacgtgaagt ccgtggaagc ctgtggctac aaaaaatggt gagccgtaga tgccgtcgga 240
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172 aatggtgaag ggagactcga agtactctga ggcttgtagg agggtaaaat agagacccag 300
173 taaaattgta ataagcagtg cttgaattat ttggtttcgg ttgttttcta ttagactatg 360
174 gtgagctcag gtgattgata ctctgatgc gagtaatacg gatgtgttta ggagtgggac 420
175 ttctagggga tttagcgggg tgatgcctgt tgggggccag tgccctccta gttggggggt 480
176 aggggctagg ctggagtggg aaaaggctca gaaaaatcct gcgaagaaaa aaacttctga 540
177 ggtaataaat aggattatcc cgtatcgaag gccttttttg acaggtggtg tgtggtggcc 600
178 ttggtatgtg ctttctcgtg ttacatcgcg ccatcattgg tatatggtta gtgtgttggg 660
W--> 179 ttantanggc ctantatgaa gaacttttgg antggaatta aatcaatngc ttggccggaa 720
W--> 180 gtcattanga nggctnaaaa ggcctgtta ngggtctggg ctnggtttta cccnaccat 780
W--> 181 ggaatncnc ccccggaacna ntgnatccct attcttaa 818
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184 <211> LENGTH: 817
185 <212> TYPE: DNA
186 <213> ORGANISM: Homo sapien
188 <220> FEATURE:
189 <221> NAME/KEY: misc_feature
190 <222> LOCATION: (1)...(817)
191 <223> OTHER INFORMATION: n = A,T,C or G
193 <400> SEQUENCE: 7
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195 cgggccctat ttcaaagatt tttaggggaa ttaattctag gacgatgggt atgaaactgt 120
196 gggttgctcc acagatttca gagcattgac cgtagtatac ccccggtcgt gtagcgggta 180
197 aagtggtttg gtttagacgt cggggaattg catctgtttt taagcctaata gtggggacag 240
W--> 198 ctcatgagtg caagacgtct tgtgatgtaa ttattatacn aatgggggct tcaatcggga 300
199 gtactactcg attgtcaacg tcaaggagtc gcaggtcgcc tggttctagg aataatgggg 360
200 gaagtatgta ggaattgaag attaatccgc cgtagtcggg gttctcctag gttcaatacc 420
201 attggtggcc aattgatttg atggttaagg gagggatcgt tgaactcgtc tgttatgtaa 480
W--> 202 aggatncctt nggatggga aggcnatnaa ggactangga tnaatggcgg gcangatatt 540
W--> 203 tcaaacngtc tctanttcct gaaacgtctg aaatgttaat aanaattaan tttngttatt 600
W--> 204 gaatnttnng gaaaagggct tacaggacta gaaaccaaata angaaaanta atnntaangg 660
W--> 205 cnttatcntn aaagtnata accnctccta tnatccacc caatngnatt cccacncnn 720
W--> 206 acnattggat nccccanttc canaaanggc cccccccgg tgnannccnc cttttgttcc 780
W--> 207 cttnantgan ggttattcnc cctngcntt atcancc 817
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210 <211> LENGTH: 799
211 <212> TYPE: DNA
212 <213> ORGANISM: Homo sapien
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215 <221> NAME/KEY: misc_feature
216 <222> LOCATION: (1)...(799)
217 <223> OTHER INFORMATION: n = A,T,C or G
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221 cataaggaga actttctgct ggcacgcgct agggacaagc gggagagcga ctccgagcgt 120
222 ctgaagcgca cgtcccagaa ggtggacttg gcaactgaaac agctgggaca catccgcgag 180
223 tacgaacagc gctgaaagt gctggagcgg gaggtccagc agtgtagccg cgtcctgggg 240
W--> 224 tgggtggcgg angcctganc cgctctgcct tgcctgcccc angtgggcgg ccacccctg 300
W--> 225 acctgcctgg gtccaaacac tgagcctgc tggcggactt caagganaac cccacangg 360
W--> 226 ggattttgct cctanantaa ggctcatctg ggccctcgcc cccccacctg gttggccttg 420

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W--> 227 tctttgangt gagcccatg tccatctggg ccaactgteng gaccaccttt ngggagtgtt      480
W--> 228 ctccctacaa ccacannatg cccggctcct cccggaaacc antcccance tgngaaggat      540
W--> 229 caagncctgn atccactnnt nctanaaccg gcncncncg cngtggaaacc cnccttntgt      600
W--> 230 tccttttctt tnagggttaa tnnegccttg gccttnccan ngtectnenc nttttccnnt      660
W--> 231 gttnaaattg ttangcnccc nccnntcccn cnnnnnnan cccgaccenn annntnnann      720
W--> 232 ncctgggggt nccnnngat tgaccenncc nccctntant tgcnttnggg nncnntgccc      780
W--> 233 ctttccctct nggganncg                                     799
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236 <211> LENGTH: 801
237 <212> TYPE: DNA
238 <213> ORGANISM: Homo sapien
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242 <222> LOCATION: (1)...(801)
243 <223> OTHER INFORMATION: n = A,T,C or G
245 <400> SEQUENCE: 9
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248 caaggacaag gccaccaggt gcgggggccc aagccacat gatccttact ctatgagcaa      180
W--> 249 aatccccctgt gggggcttct ccttgaagtc cgccancagg gctcagtctt tggaccang      240
W--> 250 caggtcatgg ggttgtnnc caactggggg ccncaacgca aaanggcnc gggcctcngn      300
W--> 251 caccatccc angacgggc tacactnctg gacctccnc tccaccaett tcatgcgtg      360
W--> 252 ttctnaccg cgnatntgtc ccantgttt cngtgcenac tccancttct nggacgtgcg      420
W--> 253 ctacatacgc ccggantcnc nctcccgtt tgccctatc cagtnccan caacaaattt      480
W--> 254 cncntantg caccnattcc cacntttnc agntttcnc nncngcttc cttntaaaag      540
W--> 255 ggttganccc cggaaaatnc cccaaagggg gggggccngg tacccaactn cccctnata      600
W--> 256 gctgaantcc catnaccnn gntcnatgg anccntccnt ttaannaen ttctnaactt      660
W--> 257 gggaaanacc ctgcncntn ccccenttaa tccnccttg cnangnnent ccccnntcc      720
W--> 258 nccnnntng gcntntnann cnaaaaaggc cnnnnaaa tctcctnnn cctcanttcg      780
W--> 259 ccancctcg aaatcgccn c                                     801
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266 <220> FEATURE:
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268 <222> LOCATION: (1)...(789)
269 <223> OTHER INFORMATION: n = A,T,C or G
271 <400> SEQUENCE: 10
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274 agatcctgcc ctacacactg gcctccctct accaccggga gaagcaggtg ttctgtccca      180
275 aataccgagg ggacactgga ggtgctagca gtgaggacag cctgatgacc agcttctgc      240
276 caggccctaa gcttgagct cccttcccta atggacacgt ggtgctgga ggcagtggcc      300
277 tgctccacc tccaccgcg ctctgcgggg cctctgctg tgatgtctcc gtacgtgtgg      360
W--> 278 tgggtgggtga gccaccgan gccagggtgg ttccgggccc gggcatctgc ctggacctcg      420
W--> 279 ccatactgga tagtgcttcc tgetgtccca ngtggcccca tccctgttta tgggctccat      480
280 tgtccagctc agccagtctg tcaactgcta tatggtgtct gccgcaggcc tgggtctggt      540
W--> 281 cccatttact ttgtacaca ggtantattt gacaagaacg anttggccaa atactcagcg      600

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Use of n and/or Xaa has been detected in the Sequence Listing.
 Review the Sequence Listing to insure a corresponding
 explanation is presented in the <220> to <223> fields of
 each sequence using n or Xaa.

VERIFICATION SUMMARY
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Input Set : A:\427d3.app.txt
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L:21 M:270 C: Current Application Number differs, Wrong Format
L:48 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:49 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:52 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:74 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:75 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:76 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:77 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:78 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:95 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:98 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:101 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:103 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:121 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:122 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:123 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:124 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:125 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
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L:153 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:155 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:179 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:180 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:181 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:202 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:203 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
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L:206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
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L:224 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
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L:226 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:227 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:228 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:229 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:230 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:231 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:232 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8

VERIFICATION SUMMARY

DATE: 12/21/2001

PATENT APPLICATION: US/10/010,940

TIME: 08:56:34

Input Set : A:\427d3.app.txt

Output Set: N:\CRF3\12212001\J010940.raw

L:233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8

L:247 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9

L:249 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9